

Serial No.: 10/806,899

IN THE SPECIFICATION

Please replace the Sequence Listing with the Substitute Sequence Listing submitted herewith. No new matter has been added.

Please replace Table 1 on page 43 with the following replacement Table 1:

TABLE 1
Primer Sequences Used for dHPLC Assay Analysis of SCN1A

Exon	Forward Primer	Reverse Primer	Size (bp)
1	CCTCTAGCTCATGGTTCATGAC (SEQ ID NO: 59)	TGCAGTAGGCAATTAGCAGC (SEQ ID NO: 60)	448
2	CTAATTAAGAAAGATCCAGTGACAG (SEQ ID NO: 61)	GCTATAAAAGTGCITACAGATCATGTAC (SEQ ID NO: 62)	356
3	CCCTGAATTGGCTAACAGCTGCAG (SEQ ID NO: 63)	CTACATTAAGACACAGTTCAAAATCC (SEQ ID NO: 64)	263
4	GGGCTACGTTCAATTGGTATG (SEQ ID NO: 65)	GCAACCTATTCTTAAGCATAAAGACTG (SEQ ID NO: 66)	358
5	AGGCTCTTGTACCTACAGC (SEQ ID NO: 67)	CATGTAGGGTCCGTCATT (SEQ ID NO: 68)	200
6	CACACGTGTTAAGCTCTCATAGT (SEQ ID NO: 69)	AGCCCCCTCAAGGTTATTATCCT (SEQ ID NO: 70)	394
7	GAACCTGACCTTCTGTTCTC (SEQ ID NO: 71)	GTTGGGCTGTTATCTTCAGTTTC (SEQ ID NO: 72)	241
8	AAAGGCCAGCAGAACGACTTG (SEQ ID NO: 73)	GGATAGAGGAACACTCAAGTCTC (SEQ ID NO: 74)	322
9	TTGAAAGTGTGAAGGCCACAC (SEQ ID NO: 75)	CCACCTGCTCTTAGGTACTC (SEQ ID NO: 76)	363
10	GCCATGCAAATACTTCAGGCC (SEQ ID NO: 77)	CACAACAGTGGTIGATTCAAGTGTG (SEQ ID NO: 78)	480
11(1)	TGAATGCTGAAATCTCCTTCTAC (SEQ ID NO: 79)	CTCAGGGTTGCTGTTGCGTCTC (SEQ ID NO: 82)	306
11(2)	GATAACGAGGCCGTAGAGAT (SEQ ID NO: 81)	TCTGTAGAAACACTGGCTGG (SEQ ID NO: 82)	315
12	CATGAAATTCACTGTGTCAAC (SEQ ID NO: 83)	CAGCTCTGAATTAGACTGTGTC (SEQ ID NO: 84)	347
13	ATCCTTGGAGGGTTAGAGT (SEQ ID NO: 85)	GCATGAAGGATGGTTGAAG (SEQ ID NO: 86)	510
14	CATTGTGGAAAATAGGATAAACG (SEQ ID NO: 87)	GCTATGCGAAACCTGTGATTG (SEQ ID NO: 88)	339
15(1)	TGAGACGGTTAGGGCAGATC (SEQ ID NO: 89)	AGAACGTCATTCAATGTGCAGC (SEQ ID NO: 90)	348
15(2)	GTCTTGGCCATCATCGTCTTC (SEQ ID NO: 91)	ACATGTGCAACAATGTGCAAGG (SEQ ID NO: 92)	350
16(1)	GTGGTGTGTTCTCTCATCAAG (SEQ ID NO: 93)	CACTGCTGCCAGTTCCTATAC (SEQ ID NO: 94)	458
16(2)	CAACAGTCTCTCATTTAGGAAAC (SEQ ID NO: 95)	ACCTTCCCACACCTATAGAATC (SEQ ID NO: 96)	353
17	CTTGGCAGGCAACCTATTAC (SEQ ID NO: 97)	CAAGCTGCACCTCCAATGAAAG (SEQ ID NO: 98)	232
18	TGGAAGCAGACACTTATCTAC (SEQ ID NO: 99)	GTGCTGTATCACCTTTCTTAATC (SEQ ID NO: 100)	234
19	CCTATTCCAATGAAATGTCAATG (SEQ ID NO: 101)	CAAGCTTACCTGAAACAGAGAC (SEQ ID NO: 102)	318
20	CTACACATGTAATGATGATTCTGT (SEQ ID NO: 103)	GCTATATACAATACTTCAGTTCT (SEQ ID NO: 104)	216
21	ACCAGAGATTACTAGGGGAAT (SEQ ID NO: 105)	CTGGGCTCATAAAACTGTACTAAC (SEQ ID NO: 106)	513
22	ACTGTCTTGGTCCAAAATCTG (SEQ ID NO: 107)	TTCGATTAAATTACCACTGTGATC (SEQ ID NO: 108)	267
23	AGCACCAGTGACATTCCAAC (SEQ ID NO: 109)	GGCAGAGAAAACACTCCAAGG (SEQ ID NO: 110)	271
24	GACACAGTTAACCAAGTTG (SEQ ID NO: 111)	TGTGAGACAAGCATGCAAGTT (SEQ ID NO: 112)	207
25	CAGGGCCAATGACTACTTGC (SEQ ID NO: 113)	CTGATTGCTGGGATGTATCTTGAATC (SEQ ID NO: 114)	477
26(1)	CAGGACTCTGAACCTTACCTTG (SEQ ID NO: 115)	ATTCACACAGATGGGTTCCCCA (SEQ ID NO: 116)	534
26(2)	TCCTGCGTTGTTAACATCGGG (SEQ ID NO: 117)	AGCGCAGCTGCAAACACTGAGAT (SEQ ID NO: 118)	504
26(3)	TGGAAGCTCAGTAAAGGGAGA (SEQ ID NO: 119)	GTAGTGTGATGGCTGATAGGGAG (SEQ ID NO: 120)	480
26(4)	CCGATGCAAACACTCAGTTCATGGGA (SEQ ID NO: 121)	TGCCCTTCTGTGCTCATGTTTCCACAC (SEQ ID NO: 122)	555
26(5)	AGAGCGATTCATGGCTTCCAATCC (SEQ ID NO: 123)	TGCTGACAAAGGGTCACTGTCT (SEQ ID NO: 124)	526

Note: Primer sequences are listed 5' to 3'. Due to the large size of exons 11, 15, 16, and 26, the exons were split into two or more overlapping amplicons.